1647

OIPE

#12

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/600,991

DATE: 09/07/2001
TIME: 14:13:11

Input Set : A:\0471-0162P.ST25.txt
Output Set: N:\CRF3\09072001\I600991.raw


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 3 <110> APPLICANT: MEDICO, Enzo
 4       MICHIELI, Paolo
 5       COLLESI, Chiara
 6       CASELLI, Gianfranco
 7       COMOGLIO, Paolo
 9 <120> TITLE OF INVENTION: RECOMBINANT PROTEINS DERIVED FROM HGF AND MSP
11 <130> FILE REFERENCE: 0471-0162P
13 <140> CURRENT APPLICATION NUMBER: US 09/600,991
C--> 14 <141> CURRENT FILING DATE: 2001-08-20
16 <160> NUMBER OF SEQ ID NOS: 22
18 <170> SOFTWARE: PatentIn version 3.1
20 <210> SEQ ID NO: 1
21 <211> LENGTH: 1725
22 <212> TYPE: DNA
23 <213> ORGANISM: Artificial Sequence
25 <220> FEATURE:
26 <223> OTHER INFORMATION: Magic F-1 DNA coding sequence
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30 ctgctcccca tcgccatccc ctatgcagag ggacaaagga aaagaagaaa tacaattcat    120
31 gaattcaaaa aatcagcaaa gactacccta atcaaaatag atccagcact gaagataaaa    180
32 accaaaaaag tgaatactgc agaccaatgt gctaatagat gtactaggaa taaaggactt    240
33 ccattcactt gcaaggcttt gtttttgat aaagcaagaa aacaatgcct ctggttcccc    300
34 ttcaatagca tgtcaagtgg agtgaaaaaa gaatttggcc atgaatttga cctctatgaa    360
35 aacaaagact acattagaaa ctgcatcatt ggtaaaggac gcagctacaa gggaacagta    420
36 tctatcacta agagtggcat caaatgtcag ccctggagtt ccatgatacc acacgaacac    480
37 agctatcggg gtaaagacct acaggaaaac tactgtcgaa atcctcgagg ggaagaaggg    540
38 ggaccctggt gtttcacaag caatccagag gtacgctacg aagtctgtga cattcctcag    600
39 tgttcagaag ttgaatgcat gacctgcaat ggggagagtt atcgaggtct catggatcat    660
40 acagaatcag gcaagatttg tcagcgctgg gatcatcaga caccacaccg gcacaaattc    720
41 ttgcctgaaa gatatcccga caagggcttt gatgataatt attgccgcaa tcccgatggc    780
42 cagccgaggc catggtgcta tactcttgac cctcacaccc gctgggagta ctgtgcaatt    840
43 aaaacatgcg ctgacaaagc ttcgggcggt ggcggttctg gtggcggtgg ctccggcggt    900
44 ggcggttctc tagagggaca aaggaaaaga agaaatacaa ttcatgaatt caaaaaatca    960
45 gcaaagacta ccctaatcaa aatagatcca gcactgaaga taaaaaccaa aaaagtgaat   1020
46 actgcagacc aatgtgctaa tagatgtact aggaataaag gacttccatt cacttgcaag   1080
47 gcttttgttt ttgataaagc aagaaaacaa tgcctctggt tccccttcaa tagcatgtca   1140
48 agtggagtga aaaaagaatt tggccatgaa tttgacctct atgaaaacaa agactacatt   1200
49 agaaactgca tcattggtaa aggacgcagc tacaagggaa cagtatctat cactaagagt   1260
50 ggcatcaaat gtcagccctg gagttccatg ataccacacg aacacagcta tcggggtaaa   1320
51 gacctacagg aaaactactg tcgaaatcct cgaggggaag aaggggggacc ctggtgtttc   1380
52 acaagcaatc cagaggtacg ctacgaagtc tgtgacattc ctcagtgttc agaagttgaa   1440
53 tgcatgacct gcaatgggga gagttatcga ggtctcatgg atcatacaga atcaggcaag   1500
54 atttgtcagc gctgggatca tcagacacca caccggcaca aattcttgcc tgaaagatat   1560
55 cccgacaagg gctttgatga taattattgc cgcaatcccg atggccagcc gaggccatgg   1620
56 tgctatactc ttgaccctca cacccgctgg gagtactgtg caattaaaac atgcgctgac   1680
57 aaagctgacg acgacgacaa acaccaccac caccaccacc actag                   1725
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Input Set.: A:\0471-0162P.ST25.txt
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<210> SEQ ID NO: 2
<211> LENGTH: 574
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Magic F-1 recombinant protein obtained combining hairpin loop and
kringle domains of human HGF and MSP

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Leu His Leu Leu Leu Leu Pro Ile Ala Ile Pro Tyr Ala Glu Gly Gln
20 25 30
Arg Lys Arg Arg Asn Thr Ile His Glu Phe Lys Lys Ser Ala Lys Thr
35 40 45
Thr Leu Ile Lys Ile Asp Pro Ala Leu Lys Ile Lys Thr Lys Lys Val
50 55 60
Asn Thr Ala Asp Gln Cys Ala Asn Arg Cys Thr Arg Asn Lys Gly Leu
65 70 75 80
Pro Phe Thr Cys Lys Ala Phe Val Phe Asp Lys Ala Arg Lys Gln Cys
85 90 95
Leu Trp Phe Pro Phe Asn Ser Met Ser Ser Gly Val Lys Lys Glu Phe
100 105 110
Gly His Glu Phe Asp Leu Tyr Glu Asn Lys Asp Tyr Ile Arg Asn Cys
115 120 125
Ile Ile Gly Lys Gly Arg Ser Tyr Lys Gly Thr Val Ser Ile Thr Lys
130 135 140
Ser Gly Ile Lys Cys Gln Pro Trp Ser Ser Met Ile Pro His Glu His
145 150 155 160
Ser Tyr Arg Gly Lys Asp Leu Gln Glu Asn Tyr Cys Arg Asn Pro Arg
165 170 175
Gly Glu Glu Gly Gly Pro Trp Cys Phe Thr Ser Asn Pro Glu Val Arg
180 185 190
Tyr Glu Val Cys Asp Ile Pro Gln Cys Ser Glu Val Glu Cys Met Thr
195 200 205
Cys Asn Gly Glu Ser Tyr Arg Gly Leu Met Asp His Thr Glu Ser Gly
210 215 220
Lys Ile Cys Gln Arg Trp Asp His Gln Thr Pro His Arg His Lys Phe
225 230 235 240
Leu Pro Glu Arg Tyr Pro Asp Lys Gly Phe Asp Asp Asn Tyr Cys Arg
245 250 255
Asn Pro Asp Gly Gln Pro Arg Pro Trp Cys Tyr Thr Leu Asp Pro His
260 265 270
Thr Arg Trp Glu Tyr Cys Ala Ile Lys Thr Cys Ala Asp Lys Ala Ser
275 280 285
Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Leu
290 295 300
Glu Gly Gln Arg Lys Arg Arg Asn Thr Ile His Glu Phe Lys Lys Ser
305 310 315 320
Ala Lys Thr Thr Leu Ile Lys Ile Asp Pro Ala Leu Lys Ile Lys Thr

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Input Set : A:\0471-0162P.ST25.txt
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134 Lys Lys Val Asn Thr Ala Asp Gln Cys Ala Asn Arg Cys Thr Arg Asn
135             340                 345                 350
137 Lys Gly Leu Pro Phe Thr Cys Lys Ala Phe Val Phe Asp Lys Ala Arg
138         355                 360                 365
140 Lys Gln Cys Leu Trp Phe Pro Phe Asn Ser Met Ser Ser Gly Val Lys
141     370                 375                 380
143 Lys Glu Phe Gly His Glu Phe Asp Leu Tyr Glu Asn Lys Asp Tyr Ile
144 385                 390                 395                 400
146 Arg Asn Cys Ile Ile Gly Lys Gly Arg Ser Tyr Lys Gly Thr Val Ser
147                 405                 410                 415
149 Ile Thr Lys Ser Gly Ile Lys Cys Gln Pro Trp Ser Ser Met Ile Pro
150             420                 425                 430
152 His Glu His Ser Tyr Arg Gly Lys Asp Leu Gln Glu Asn Tyr Cys Arg
153         435                 440                 445
155 Asn Pro Arg Gly Glu Glu Gly Gly Pro Trp Cys Phe Thr Ser Asn Pro
156     450                 455                 460
158 Glu Val Arg Tyr Glu Val Cys Asp Ile Pro Gln Cys Ser Glu Val Glu
159 465                 470                 475                 480
161 Cys Met Thr Cys Asn Gly Glu Ser Tyr Arg Gly Leu Met Asp His Thr
162                 485                 490                 495
164 Glu Ser Gly Lys Ile Cys Gln Arg Trp Asp His Gln Thr Pro His Arg
165             500                 505                 510
167 His Lys Phe Leu Pro Glu Arg Tyr Pro Asp Lys Gly Phe Asp Asp Asn
168         515                 520                 525
170 Tyr Cys Arg Asn Pro Asp Gly Gln Pro Arg Pro Trp Cys Tyr Thr Leu
171     530                 535                 540
173 Asp Pro His Thr Arg Trp Glu Tyr Cys Ala Ile Lys Thr Cys Ala Asp
174 545                 550                 555                 560
176 Lys Ala Asp Asp Asp Asp Lys His His His His His His His
177                 565                 570
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181 <211> LENGTH: 1692
182 <212> TYPE: DNA
183 <213> ORGANISM: Artificial Sequence
185 <220> FEATURE:
186 <223> OTHER INFORMATION: Metron F-1 DNA coding sequence
188 <400> SEQUENCE: 3
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190 tcgccattga atgacttcca agtgctccgg ggcacagagc tacagcacct gctacatgcg    120
191 gtggtgcccg ggccttggca ggaggatgtg gcagatgctg aagagtgtgc tggtcgctgt    180
192 gggcccttaa tggactgccg ggccttccac tacaacgtga gcagccatgg ttgccaactg    240
193 ctgccatgga ctcaacactc gccccacacg aggctgcggc gttctgggcg ctgtgacctc    300
194 ttccagaaga aagactacgt acggacctgc atcatgaaca atggggttgg gtaccggggc    360
195 accatggcca cgaccgtggg tggcctgccc tgccaggctt ggagccacaa gttcccgaat    420
196 gatcacaagt acacgcccac tctccggaat ggcctggaag agaacttctg ccgtaaccct    480
197 gatggcgacc ccggaggtcc ttggtgctac acaacagacc ctgctgtgcg cttccagagc    540
198 tgcggcatca aatcctgccg ggaggccgcg tgtgtctggt gcaatggcga ggaataccgc    600
199 ggcgcggtag accgcacgga gtcagggcgc gagtgccagc gctgggatct tcagcacccg    660
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PATENT APPLICATION: US/09/600,991 TIME: 14:13:11

Input Set : A:\0471-0162P.ST25.txt
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200 caccagcacc ccttcgagcc gggcaagttc ctcgaccaag gtctggacga caactattgc 720
201 cggaatcctg acggctccga gcggccatgg tgctacacta cggatccgca gatcgagcga 780
202 gagttctgtg acctcccccg ctgcgggtcc gaggcacagc cccgcctcga gggcggtggc 840
203 ggttctggtg gcggtggctc cggcggtggc ggttctctag agggacaaag gaaaagaaga 900
204 aatacaattc atgaattcaa aaaatcagca aagactaccc taatcaaaat agatccagca 960
205 ctgaagataa aaaccaaaaa agtgaatact gcagaccaat gtgctaatag atgtactagg 1020
206 aataaaggac ttccattcac ttgcaaggct tttgtttttg ataaagcaag aaaacaatgc 1080
207 ctctggttcc ccttcaatag catgtcaagt ggagtgaaaa aagaatttgg ccatgaattt 1140
208 gacctctatg aaaacaaaga ctacattaga aactgcatca ttggtaaagg acgcagctac 1200
209 aagggaacag tatctatcac taagagtggc atcaaatgtc agccctggag ttccatgata 1260
210 ccacacgaac acagctatcg gggtaaagac ctacaggaaa actactgtcg aaatcctcga 1320
211 ggggaagaag ggggaccctg gtgtttcaca agcaatccag aggtacgcta cgaagtctgt 1380
212 gacattcctc agtgttcaga agttgaatgc atgacctgca atggggagag ttatcgaggt 1440
213 ctcatggatc atacagaatc aggcaagatt tgtcagcgct gggatcatca gacaccacac 1500
214 cggcacaaat tcttgcctga aagatatccc gacaagggct ttgatgataa ttattgccgc 1560
215 aatcccgatg gccagccgag gccatggtgc tatactcttg accctcacac ccgctgggag 1620
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217 caccaccact ag 1692
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222 <212> TYPE: PRT
223 <213> ORGANISM: Artificial Sequence
225 <220> FEATURE:
226 <223> OTHER INFORMATION: Metron F-1 recombinant protein obtained combining hairpin loop and
227 kringle domains of human HGF and MSP
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232 1 5 10 15
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235 20 25 30
237 Glu Leu Gln His Leu Leu His Ala Val Val Pro Gly Pro Trp Gln Glu
238 35 40 45
240 Asp Val Ala Asp Ala Glu Glu Cys Ala Gly Arg Cys Gly Pro Leu Met
241 50 55 60
243 Asp Cys Arg Ala Phe His Tyr Asn Val Ser Ser His Gly Cys Gln Leu
244 65 70 75 80
246 Leu Pro Trp Thr Gln His Ser Pro His Thr Arg Leu Arg Arg Ser Gly
247 85 90 95
249 Arg Cys Asp Leu Phe Gln Lys Lys Asp Tyr Val Arg Thr Cys Ile Met
250 100 105 110
252 Asn Asn Gly Val Gly Tyr Arg Gly Thr Met Ala Thr Thr Val Gly Gly
253 115 120 125
255 Leu Pro Cys Gln Ala Trp Ser His Lys Phe Pro Asn Asp His Lys Tyr
256 130 135 140
258 Thr Pro Thr Leu Arg Asn Gly Leu Glu Glu Asn Phe Cys Arg Asn Pro
259 145 150 155 160
261 Asp Gly Asp Pro Gly Gly Pro Trp Cys Tyr Thr Thr Asp Pro Ala Val
262 165 170 175
264 Arg Phe Gln Ser Cys Gly Ile Lys Ser Cys Arg Glu Ala Ala Cys Val

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Input Set : A:\0471-0162P.ST25.txt
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265                 180                 185                 190
267 Trp Cys Asn Gly Glu Glu Tyr Arg Gly Ala Val Asp Arg Thr Glu Ser
268         195                 200                 205
270 Gly Arg Glu Cys Gln Arg Trp Asp Leu Gln His Pro His Gln His Pro
271     210                 215                 220
273 Phe Glu Pro Gly Lys Phe Leu Asp Gln Gly Leu Asp Asp Asn Tyr Cys
274 225                 230                 235                 240
276 Arg Asn Pro Asp Gly Ser Glu Arg Pro Trp Cys Tyr Thr Thr Asp Pro
277                 245                 250                 255
279 Gln Ile Glu Arg Glu Phe Cys Asp Leu Pro Arg Cys Gly Ser Glu Ala
280                 260                 265                 270
282 Gln Pro Arg Leu Glu Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly
283         275                 280                 285
285 Gly Gly Gly Ser Leu Glu Gly Gln Arg Lys Arg Arg Asn Thr Ile His
286     290                 295                 300
288 Glu Phe Lys Lys Ser Ala Lys Thr Thr Leu Ile Lys Ile Asp Pro Ala
289 305                 310                 315                 320
291 Leu Lys Ile Lys Thr Lys Lys Val Asn Thr Ala Asp Gln Cys Ala Asn
292                 325                 330                 335
294 Arg Cys Thr Arg Asn Lys Gly Leu Pro Phe Thr Cys Lys Ala Phe Val
295                 340                 345                 350
297 Phe Asp Lys Ala Arg Lys Gln Cys Leu Trp Phe Pro Phe Asn Ser Met
298         355                 360                 365
300 Ser Ser Gly Val Lys Lys Glu Phe Gly His Glu Phe Asp Leu Tyr Glu
301     370                 375                 380
303 Asn Lys Asp Tyr Ile Arg Asn Cys Ile Ile Gly Lys Gly Arg Ser Tyr
304 385                 390                 395                 400
306 Lys Gly Thr Val Ser Ile Thr Lys Ser Gly Ile Lys Cys Gln Pro Trp
307                 405                 410                 415
309 Ser Ser Met Ile Pro His Glu His Ser Tyr Arg Gly Lys Asp Leu Gln
310                 420                 425                 430
312 Glu Asn Tyr Cys Arg Asn Pro Arg Gly Glu Glu Gly Gly Pro Trp Cys
313         435                 440                 445
315 Phe Thr Ser Asn Pro Glu Val Arg Tyr Glu Val Cys Asp Ile Pro Gln
316     450                 455                 460
318 Cys Ser Glu Val Glu Cys Met Thr Cys Asn Gly Glu Ser Tyr Arg Gly
319 465                 470                 475                 480
321 Leu Met Asp His Thr Glu Ser Gly Lys Ile Cys Gln Arg Trp Asp His
322                 485                 490                 495
324 Gln Thr Pro His Arg His Lys Phe Leu Pro Glu Arg Tyr Pro Asp Lys
325                 500                 505                 510
327 Gly Phe Asp Asp Asn Tyr Cys Arg Asn Pro Asp Gly Gln Pro Arg Pro
328         515                 520                 525
330 Trp Cys Tyr Thr Leu Asp Pro His Thr Arg Trp Glu Tyr Cys Ala Ile
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340 <210> SEQ ID NO: 5
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VERIFICATION SUMMARY DATE: 09/07/2001
PATENT APPLICATION: US/09/600,991 TIME: 14:13:12

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